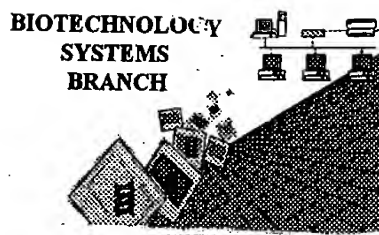


1600 1/2



## RAW SEQUENCE LISTING ERROR REPORT

TECH CENTER 1600/2900

DEC 8 0 2002

#10

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/869,185A  
Source: Q1PE  
Date Processed by STIC: 12/12/02

**THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.**

**PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:**

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)

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**TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:**

**<http://www.uspto.gov/web/offices/pac/checker>**

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:  
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202  
Or  
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

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Does Not Comply  
Corrected Diskette Needed

OIPE

## RAW SEQUENCE LISTING

DATE: 12/12/2002

PATENT APPLICATION: US/09/869,185A

TIME: 14:45:26

Input Set : A:\EP.txt

Output Set: N:\CRF4\12122002\I869185A.raw

3 <110> APPLICANT: Ashikari, Toshihiko  
4 Ochiai, Misa  
6 <120> TITLE OF INVENTION: Method of Breeding Yeast  
8 <130> FILE REFERENCE: 46221  
10 <140> CURRENT APPLICATION NUMBER: US 09/869,185A  
12 <141> CURRENT FILING DATE: 2001-06-25  
14 <150> PRIOR APPLICATION NUMBER: PCT/JP00/07491  
16 <151> PRIOR FILING DATE: 2000-10-26  
E--> 18 <160> NUMBER OF SEQ ID NOS: (28) - counted 30

## ERRORED SEQUENCES

496 <210> SEQ ID NO: 30  
498 <211> LENGTH: 30  
500 <212> TYPE: DNA  
502 <213> ORGANISM: Artificial Sequence  
504 <220> FEATURE:  
506 <223> OTHER INFORMATION: Sequence of synthetic DNA to prepare a combined FRT103-a /  
FRT3-a sequence  
E--> 508 <400> SEQUENCE: (29) - change to 30  
509 agctttccta tacttttctag agaataggag 30

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/869,185A

DATE: 12/12/2002  
TIME: 14:45:28

Input Set : A:\EP.txt

Output Set: N:\CRF4\12122002\I869185A.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:3; Line(s) 63  
Seq#:4; Line(s) 80  
Seq#:6; Line(s) 114  
Seq#:7; Line(s) 131  
Seq#:9; Line(s) 165  
Seq#:10; Line(s) 182  
Seq#:11; Line(s) 199  
Seq#:12; Line(s) 216  
Seq#:29; Line(s) 490  
Seq#:30; Line(s) 506

## VERIFICATION SUMMARY

DATE: 12/12/2002

PATENT APPLICATION: US/09/869,185A

TIME: 14:45:28

Input Set : A:\EP.txt

Output Set: N:\CRF4\12122002\I869185A.raw

L:508 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:30 differs:29  
L:18 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (28) Counted (30)